

1 CACAGACAGGCTCTTAACTGTTTATATCACTCACTTGGGAAAGTGTCTCAAGCTGTTCTA  
61 CAAATCCATGCAAAGGCCGTTTAAAAATAGCAGCGAAGGTCCTGGACTCGGTCTCGTCCA  
121 GCACAGCCCCTTGGCTCTCTCTCTGGGCTCTGGCCGCCTGGCCCCCGGGGACCCACACGA  
181 GGTCATGGCGTGCTTCGGGCAGGGGGCGGGGATCCCATAGACACCTCAGCTCCTTAAGA  
241 GTTCTCCGCCTGGGCCAGGACGAGCATGGGGGTCCCCACTGATGCCCCGAGACAGTGCCCC  
301 TGTGTGTGTGAGCCCTCGACCCACATAACAGAGAGGTGTCCTGATGCCCTCTGTCCTCTC  
361 CAGGTGGATCTAGGATCCGGCTTCCAACATGTGGCAGCTCTGGGCCTCCCTCTGCTGCCT  
421 GCTGGTGTGAGCCAATGCCCGGAGCAGGCCCTCTTTCCATCCCGTGTCGGATGAGCTGGT  
481 CAACTATGTCAACAAACGGAATACCACGTGGCAGGCCGGGCACAACTTCTACAACGTGGA  
541 CATGAGCTACTTGAAGAGGCTATGTGGTACCTTCCTGGGTGGGCCCCAAGCCACCCCAGAG  
601 AGTTATGTTTACCGAGGACCTGAAGCTGCCTGCAAGCTTCGATGCACGGGAACAATGGCC  
661 ACAGTGTCCCACCATCAAAGAGATCAGAGACCAGGGCTCCTGTGGCTCCTGCTGGGTAAG  
721 GCCCTGCTGGCTGGTGGGGAAGCGCTGGAGAGAAAGTGGGAGCAACACTGGAGAGTCTTG  
781 GGGGATTCCGGGTGGGGACAACCTCTGACAAGGCAAGTTATAGAACTTTCTGAGTCCCAG  
781 TTTCCATCAGTACAAAAATCACAATCCCTCTGGCCATGAATGATGGCGAGGATTAGGTGG  
841 AGTGGCGGGCAGAGCATCCAGCAGATTGCAAGTCCACGTGTACAGGTGGCGAAGCAGCTC  
901 CCTTTCCCTGACATGCTGGCCCGTCCGCAAATACCAGGAGCTCTCACTGCTACTCTGCTT  
961 CAAGAAAGCATCCCTTTAGTGTGAGTGTCTTAATTTTGTCAATTAATTGTGGTAA  
1021 AATACACGTAACAGAAATGTAATAATCTTAGCAATCTTCTTTTGTTTTCTTTTTCTTTT  
1081 TTTTTTTTTTTTTTTT

FIGURE 1

1 MWQLWASLCCLLVLANARSRPSFHPVSELVNYVKNRNTTWQAGHNFYNVDMSYLRKLCG  
61 TFLGGPKPPQVRMFTEDLKLPASFDAREQWPQCPTIKEIRDQSCGSCWVRPCWLVGKRW  
121 RESGSNTGESWGIRGGDNSDKASYRNFLSPSFHQYKNHNPSGHE

**FIGURE 2**

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1 TCCAAGTCCTGGATGAGCTCGGAAGACACGTGCTGCTGCGGAAGGACTGTGGCCCAGTGG  
61 ACACCAAGGTTCCAGGTGCTGGGGAGCCCAAGTCAGCCTTCACTCAGGGCTCAGCCATGA  
121 TTTCTTCTCTGTCCCAAACCATCCAGACAACAGAAACGAGACTTTTCAGCTCAGTCATTT  
181 CCCTGTTGAATGAGGATCCCCTGTCCCAGGACTTGCCTGTGAAGATGGCTTCAATCTTCA  
241 AGAACTTTGTCAATTACCTATAACCGGACATATGAGTCAAAGGAAGAAGCCCGGTGGCGCC  
301 TGTCCGTCTTTGTCAATAACATGGTGCGAGCACAGAAGATCCAGGCCCTGGACCGTGGCA  
361 CAGCTCAGTATGGAGTCACCAAGTTCAGTGATCTCACAGAGGAGGAGTTCGCACTATCT  
421 ACCTGAATACTCTCCTGAGGAAAGAGCCTGGCAACAAGATGAAGCAAGCCAAGTCTGTGG  
481 GTGACCTCGCCCCACCTGAATGGGACTGGAGGAGTAAGGGGGCTGTACAAAAGTCAAAG  
541 ACCAGGGCATGTGTGGCTCCTGCTGGGCCTTCTCAGTCACAGGCAATGTGGAGGGCCAGT  
601 GGTTCCTCAACCAGGGGACCCTGCTCTCCCTCTCTGAACAGGAGCTCTTGGACTGTGACA  
661 AGATGGACAAGGCCTGCATGGGCGGCTTGCCCTCCAATGCCTACTCGGCCATAAAGAATT  
721 TGGGAGGGCTGGAGACAGAGGATGACTACAGCTACCAGGGTCACATGCAGTCCTGCAACT  
781 TCTCAGCAGAGAAGGCCAAGGTCTACATCAATGACTCCGTGGAGCTGAGCCAGAACGAGC  
841 AGAAGCTGGCAGCCTGGCTGGCCAAGAGAGGCCCAATCTCCGTGGCCATCAATGCCTTTG  
901 GCATGCAGTTTTACCGCCACGGGATCTCCCGCCCTCTCCGGCCCCTCTGCAGCCCTTGGC  
961 TCATTGACCATGCGGTGTTGCTTGTGGGCTACGGCAACCGCTCTGACGTTCCCTTTTGGG  
1021 CCATCAAGAACAGCTGGGGCACTGACTGGGGTGAGAAGGGTTACTACTACTTGCATCGCG  
1081 GGTCCGGGGCCTGTGGCGTGAACACCATGGCCAGCTCGGCGGTGGTGGACTGAAGAGGGG  
1141 CCCCCAGCTCGGGACCTGGTGCTGATCAGAGTGGCTGCTGCCCCAGCCTGACATGTGTCC  
1201 AGGCCCCTCCCCGGGAGGTACAGCTGGCAGAGGGAAAGGCACTGGGTACCTCAGGGTGAG  
1261 CAGAGGGCACTGGGCTGGGGCACAGCCCCTGCTT

FIGURE 3

1 MISSLSQNHDPNRNETFSSVISLLNEDPLSQDLPVKMASIFKNFVITYNRTYESKEEARW  
61 RLSVFVNNMVRAQKIQALDRGTAQYGVTKFSDLTEEEFRTIYLNTLLRKEPGNKMKAQS  
121 VGD LAPPEWDWRSKGAVTKVKDQGMCGSCWAFSVTGNVEGQWFLNQGTLLSLSEQELLDC  
181 DKMDKACMGGLPSNAYSIAKNLGGLETEDDYSYQGHMQSCNFSAEKAKVYINDSVELSON  
241 EQKLAAWLAKRGPISVAINAFGMQFYRHGISRPLRPLCSPWLIDHAVLLVGYGNRSDVPF  
301 WAIKNSWGTDWGEKGYYYLHRGSGACGVNTMASSAVVD

**FIGURE 4**

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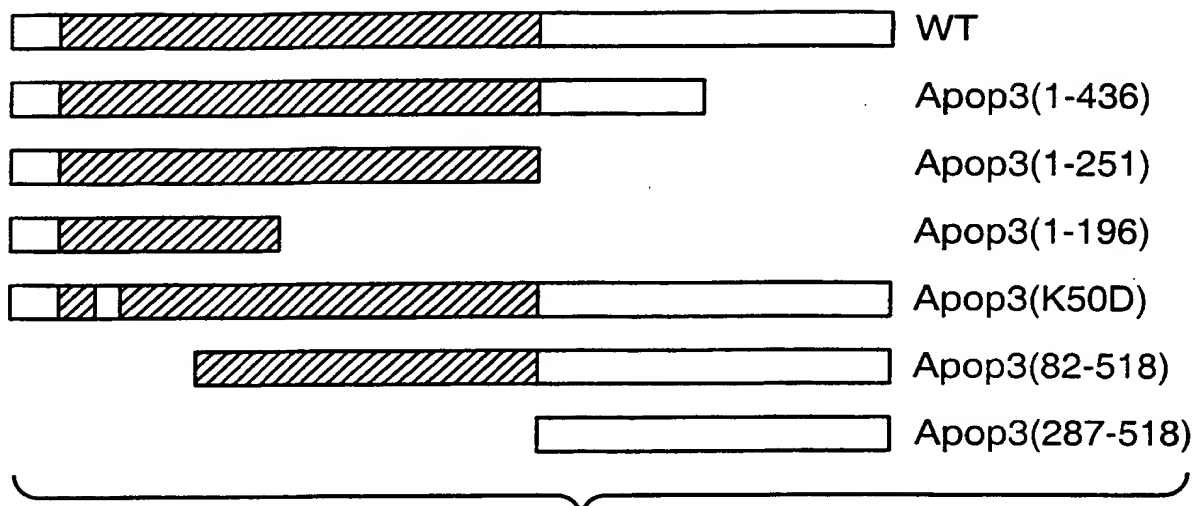
1 atgtcgtgcg tcaagttatg gccagcggt gccccgccc ccttggtgtc catcgaggaa  
61 ctggagaacc aggagctcgt cggcaaagac gggttcggca cagtgttcg ggcgcaacat  
121 aggaagtggg gctacgatgt ggcggtcaag atcgtaaact cgaaggcgat atccaggag  
181 gtcaaggcca tggcaagtct ggataacgaa ttcgtgttgc gcctagaagg ggttatcgag  
241 aaggtgaact gggaccaaga tcccaagccg gctctggtga ctaaattcat ggagaacggc  
301 tccttgctcg ggctgctgca gtcccagtg cctcggccct ggccgctcct ttgccgctg  
361 ctgaaagaag tgggtgcttg gatgttttac ctgcacgacc agaaccggg gtcctgcac  
421 cgggacctca agccatccaa cgtcctgccg gaccagagc tgcacgtcaa gctggcagat  
481 tttggcctgt ccacatttca gggaggctca cagtcaggga cagggtccg ggagccagg  
541 ggcaccctgg gctacttggc ccagaaactg tttgttaacg taaaccggaa ggcctccaca  
601 gccagtgcg tctacagctt cgggatccta atgtgggcag tgcttgctgg aagagaagtt  
661 gagttgccaa ccgaaccatc actcgtgtac gaagcagtgt gcaacaggca gaaccggcct  
721 tcattggctg agctgcccc a gccgggcct gagactccg gcttagaagg actgaaggag  
781 ctaatgcagc tctgctggag cagtgcgccc aaggacagac cctccttcca ggaatgccta  
841 ccaaaaactg atgaagtctt ccagatggtg gagaacaata tgaatgctgc tgtctccacg  
901 gtaaaggatt tcctgtctca gctcaagagc agcaatagga gatttttctat cccagagtca  
961 ggccaaggag ggacagaaat ggatggcttt aggagaacca tagaaaacca gcactctcgt  
1021 aatgatgtca tgggtttctga gtggctaaac aaactgaatc tagaggagcc tcccagctct  
1081 gttcctaaaa aatgcccag cttaccaag aggagcaggg cacaagagga gcaggttcca  
1141 caagcctgga cagcaggcac atcttcagat tcgatggccc aacctccca gactccagag  
1201 acctcaactt tcagaaacca gatgcccagc cctacctcaa ctggaacacc aagtccctgga  
1261 ccccagggga atcagggggc tgagagacaa ggcatgaact ggtcctgcag gaccccgag  
1321 ccaaatccag taacaggggc accgctcgtt aacatataca actgctctgg ggtgcaagtt  
1381 ggagacaaca actacttgac tatgcaacag acaactgcct tgcccacatg gggcttggca  
1441 ccttcgggca aggggagggg cttgcagcac ccccaccag taggttcgca agaaggccct  
1501 aaagatcctg aagcctggag caggccacag ggttggtata atcatagcg gaaataa

FIGURE 5

1 MSCVKLWPSGAPAPLVSIEELENQELVGKDGFGTVFRAQHRKWGYDVAVKIVNSKAISRE  
61 VKAMASLDNEFVLRLEGVIEKVNWDQDPKALVTKFMENGSLSGLLQSQCPRPWPLLCLRL  
121 LKEVVLGMFYLDQNPVLLHRDLKPSNVLPDPELVKLAADFGLSTFQGGSQSGTGSGEPG  
181 GTLGYLAPELFVNVNRKASTASDVYSFGILMWAVLAGREVELPTEPSLVYEAVCNRQNRP  
241 SLAELPQAGPETPGLEGLKELMQLCWSSEPDKRPSFQECLPKTDEVFQMVENNMNAAVST  
301 VKDFLSQLKSSNRRFSIPESGQGGTEMDGFRRTIENQHSRNDVMVSEWLNKLNLEPPSS  
361 VPKKCPSLTKRSRAQEEQVPQAWTAGTSSDSMAQPPQTPETSTFRNQMPSPSTSTGTPSPG  
421 PRGNQGAERQGMNWSCRTPEPNPVTGRPLVNIYNCSGVQVGDNNYLTMQQTALPTWGLA  
481 PSGKGRGLQHPPVGSQEGPKDPEAWSRPQGWYNHSGKZ

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## FIGURE 6

**FIG.\_7**

		Binding To RIP
<b>FIG._8</b>	Apop3	++++
	Apop3(1-436)	+
	Apop3(1-251)	-
	Apop3(1-196)	-
	Apop3(K50D)	++++
	Apop3(82-518)	+++
	Apop3(287-518)	-

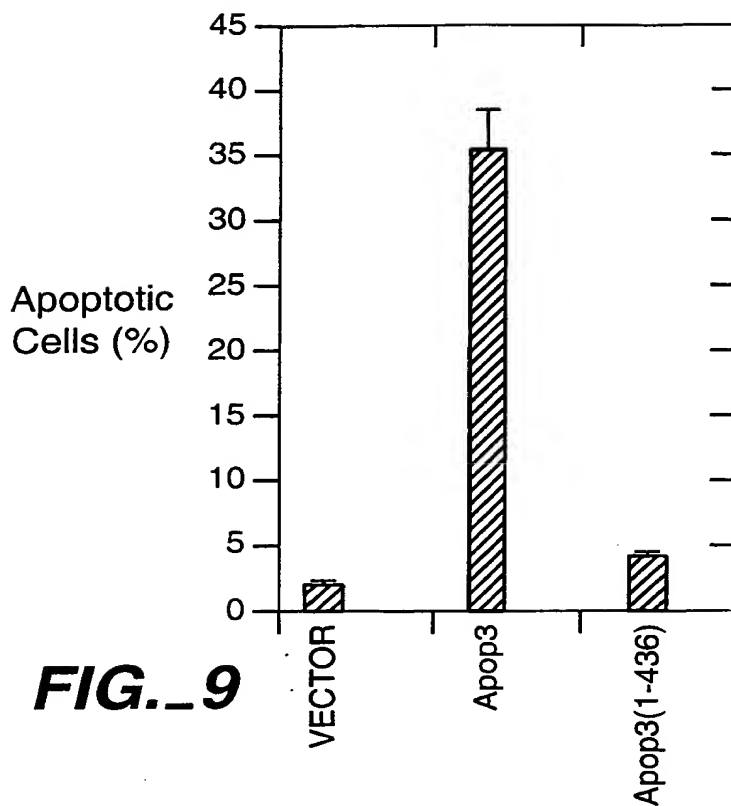


FIG.\_9

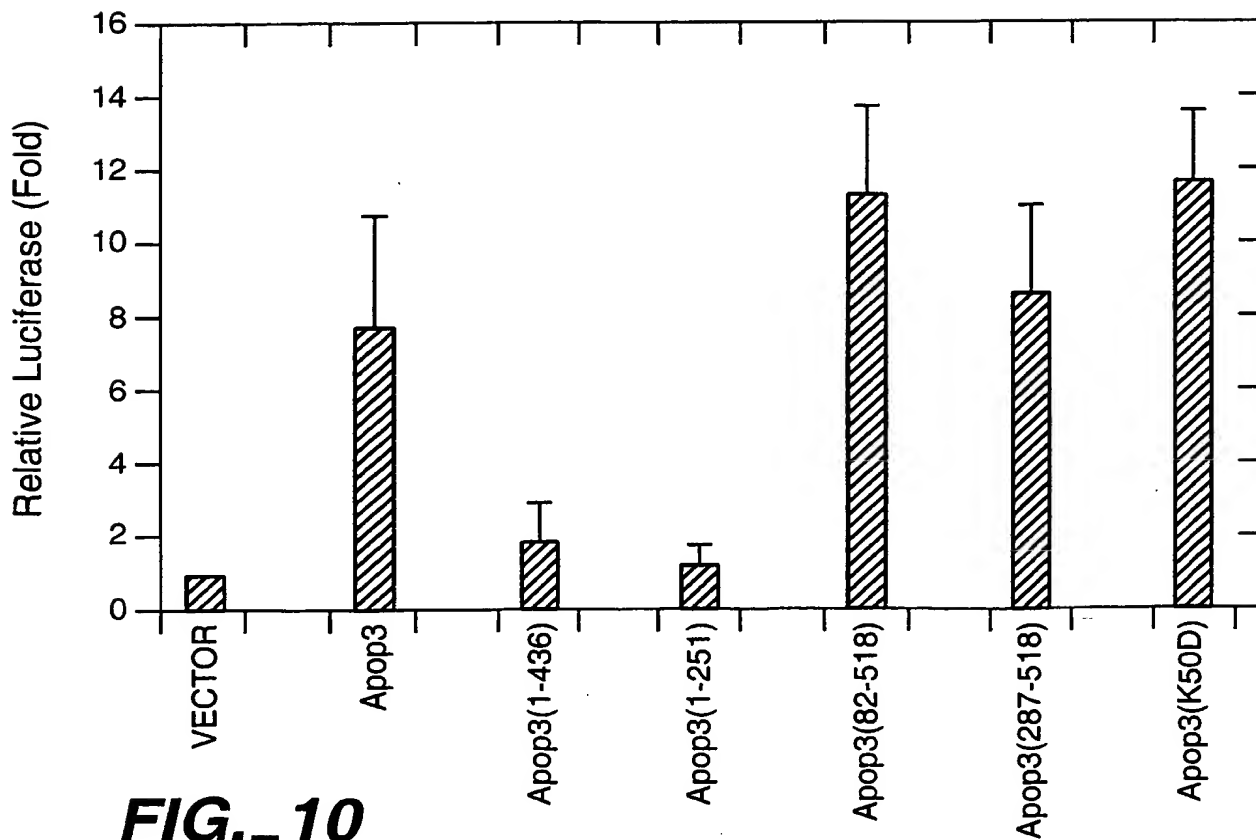


FIG.\_10